

GenCore version 4.5
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OM nucleic - nucleic search, using sw model

Run on: August 15, 2002, 08:29:09 ; Search time 211.29 Seconds
(without alignments)
2681.536 Million cell updates/sec

Title: US-09-622-613a-14

Perfect score: 330
Sequence: 1 cagaactggcgctacttcca.....ctggtatcggtcggtccgc 330

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 1736436 seqs, 858457221 residues

Total number of hits satisfying chosen parameters: 3472872

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : N_Geneseq_032802.*
1: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1980.DAT.*
2: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1981.DAT.*
3: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA1982.DAT.*
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21: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2000.DAT.*
22: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2001A.DAT.*
23: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2001B.DAT.*
24: /SIDSL1/gcgdata/geneseq/geneseqn-emb1/NA2002.DAT.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	330	100.0	330	AAZ08130	Rana catesbeiana o
2	330	100.0	333	AAZ08131	Recombinant Met(-1
3	327	99.1	330	AAZ08134	Recombinant RacOR1
4	327	99.1	333	AAZ08135	Recombinant Met(-1
5	326.8	99.0	330	AAZ08132	Recombinant RacOR1
6	326.8	99.0	333	AAZ08133	Recombinant Met(-1
7	95	28.8	95	AAZ08144	PCR primer-3 for s
8	94.4	28.6	96	AAZ08148	PCR primer-2 for a
9	92.8	28.1	96	AAZ08141	PCR primer-1 for s

10	91.2	27.6	96	AAZ08147	PCR primer-1 for a
11	85	25.8	97	AAZ08139	PCR primer-1 for s
12	71.8	21.8	86	AAZ08140	PCR primer-2 for s
13	50	15.2	318	AAZ19767	Recombinant frog O
14	39.2	11.9	19205	AAZ34685	Human DNA for a no
15	39	11.8	436	ABAA4186	Human breast cell
16	39	11.8	436	ABAA5635	Human foetal liver
17	39	11.8	436	ABAA2419	Human brain expres
18	39	11.8	436	AAK02925	Human bone marrow
19	39	11.8	436	AAK28369	Probe #2865 for ge
20	39	11.8	436	AAI12933	Probe #2862 used t
21	39	11.8	436	AAI34296	R. pipiens recombi
22	39	11.8	436	AAI02854	R. pipiens recombi
23	38.2	11.6	315	AAAT94959	R. pipiens recombi
24	38.2	11.6	318	AAAT94958	R. pipiens recombi
25	38.2	11.6	321	AAAT94954	R. pipiens recombi
26	38.2	11.6	333	AAAT94957	R. pipiens recombi
27	38.2	11.6	336	AAAT94955	R. pipiens recombi
28	38.2	11.6	753	AAAT94972	R. pipiens recombi
29	38.2	11.6	768	AAAT94973	R. pipiens recombi
30	38.2	11.6	1065	AAAT94971	R. pipiens recombi
31	38.2	11.6	1065	AAAT94963	R. pipiens recombi
32	38.2	11.6	1065	AAAT94967	R. pipiens recombi
33	38.2	11.6	1074	AAAT94968	R. pipiens recombi
34	38.2	11.6	1098	AAAT94970	R. pipiens recombi
35	38.2	11.6	1137	AAAT94964	R. pipiens recombi
36	38	11.5	5997	AAO12188	Odonotoglossum ring
37	38	11.5	6597	AAO38106	ORSV CDNA, Odonot
38	37.6	11.4	312	AAZ08124	Rana pipiens liver
39	37.6	11.4	312	AAZ08125	Recombinant RapLR1
40	37.6	11.4	312	AAZ08128	Recombinant RapLR1
41	37.6	11.4	315	AAZ08126	Recombinant Met(-1
42	37.6	11.4	315	AAZ08127	Recombinant Met(-1
43	37.6	11.4	315	AAZ08129	Recombinant Met(-1
44	37.6	11.4	2855	AAZ08136	Rana pipiens ribom
45	37.4	11.3	2574	AAV26293	Recombinant botuli

ALIGNMENTS

RESULT 1	
AAZ08130	standard; CDNA: 330 BP.
ID	AAZ08130
XX	AAZ08130;
AC	25-JAN-2000 (first entry)
XX	
DT	Rana catesbeiana oocyte ribonuclease (RacOR1) encoding cDNA.
XX	
DE	
XX	Rana catesbeiana oocyte ribonuclease; RacOR1; covalently bound; CD22;
XX	IL2 antibody; ligand binding moiety; cancerous B cell; kaposi's Sarcoma;
KW	human chorionic gonadotropin; hcg; recombinant ribonuclease; bullfrog;
KW	signal peptide; cytotoxic fusion protein; cancer; autoimmune disease;
KW	RNase; ss.
XX	
OS	Rana catesbeiana.
OS	Synthetic.
XX	
FH	Key
FT	mat_peptide
FT	Location/Qualifiers
FT	1..330
FT	/*tag= a
FT	/product= "RacOR1"
FT	/note= "Rana catesbeiana oocyte ribonuclease"
XX	
PN	W09950398-A2.
XX	
XX	07-OCT-1999.
PD	
XX	
XX	26-MAR-1999; 99WO-US06641.
PF	
XX	
XX	27-MAR-1998; 98US-0079751.

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XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
PA Newton DL, Rybak SM.
XX
XX WPI; 1999-610847/52.
XX P-PSDB; AAY28872.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX Claim 20; Page 62; 71pp; English.
XX
XX The present sequence is a synthetic cDNA encoding Rana catesbeiana oocyte
XX ribonuclease (RacOR1), modified for expression in E. coli. Carboxy
XX terminal end of RacOR1 has a covalently bound ligand binding moiety,
XX which can be a IL2 antibody directed against CD22 on cancerous B cells or
XX human chorionic gonadotropin (hCG) effective against Kaposi's Sarcoma
XX cells. Recombinant ribonucleases can be expressed in bacteria without an
XX N-terminal methionine due to the presence of a signal peptide that is
XX cleaved by bacteria. The soluble expression of ribonuclease allows the
XX proteins to be fused in-frame with ligand binding moieties to form
XX cytotoxic fusion proteins. They can be used for treatment of cancer and
XX autoimmune diseases.
XX
XX Sequence 330 BP; 82 A; 95 C; 56 G; 97 T; 0 other:
XX
Query Match          100.0%; Score 330; DB 20; Length 330;
Best Local Similarity 100.0%; Pred. No. 4.4e-92;
Matches 330; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 cagaactgggtactcttcagcagaacacatcatcaaacctcgcgactcgtgaacact 60
DB 1 cagaactgggtactcttcagcagaacacatcatcaaacctcgcgactcgtgaacact 60
QY 61 atcattggacacacacatctacatcgtgtggtgcagtcgcaacagtggttaacatttc 120
DB 61 atcattggacacacacatctacatcgtgtggtgcagtcgcaacagtggttaacatttc 120
QY 121 atctctctgctactactgtttaagatcatctgactgtgtgtttaacaagaagttctg 180
DB 121 atctctctgctactactgtttaagatcatctgactgtgtgtttaacaagaagttctg 180
QY 121 atctctctgctactactgtttaagatcatctgactgtgtgtttaacaagaagttctg 180
DB 121 atctctctgctactactgtttaagatcatctgactgtgtgtttaacaagaagttctg 180
QY 181 tctactactcgtttccagctgaacactgtgactcgtactctatcaactccgctcgtgc 240
DB 181 tctactactcgtttccagctgaacactgtgactcgtactctatcaactccgctcgtgc 240
QY 241 cgtactctctcgtactcgtgaactaactacatctgcgtttaaagcgaaaacagttaccg 300
DB 241 cgtactctctcgtactcgtgaactaactacatctgcgtttaaagcgaaaacagttaccg 300
QY 301 gtccattcgtcgtgtatcgcgttgcccg 330
DB 301 gtccattcgtcgtgtatcgcgttgcccg 330
XX
RESULT 2
ID AAZ08131 standard; cDNA; 333 BP.
XX
XX AAZ08131;
XX
XX 25-JAN-2000 (first entry)
XX
XX Recombinant Met(-1) RacOR1 encoding cDNA.
XX
XX Recombinant Met(-1) Rana catesbeiana oocyte ribonuclease; RacOR1; CD22;
XX RNase; covalently bound; IL2 antibody; ligand binding moiety;
XX cancerous B cell; Kaposi's sarcoma; human chorionic gonadotropin; hCG;
XX signal peptide; recombinant ribonuclease; cytotoxic fusion protein;
XX cancer; bullfrog; autoimmune disease; ss.
XX

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OS Rana catesbeiana.
OS Synthetic.
XX
XX Key Location/Qualifiers
XX mat_peptide 1..333
XX FT /*tag- a
XX FT /*product= "Recombinant Met(-1) RacOR1"
XX FT /*note= "Rana catesbeiana oocyte ribonuclease"
XX FT 1..3
XX FT /*tag- b
XX FT /*note= "Additional ATG codon not found in RacOR1"
XX
XX W09950398-A2.
XX
XX PD 07-OCT-1999.
XX
XX PF 26-MAR-1999; 99MO-US06641.
XX
XX PR 27-MAR-1998; 98US-0079751.
XX
XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
XX
XX Newton DL, Rybak SM.
XX
XX WPI; 1999-610847/52.
XX P-PSDB; AAY28873.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX Disclosure; Page 63; 71pp; English.
XX
XX The present sequence is a cDNA encoding recombinant Rana catesbeiana
XX oocyte ribonuclease (RacOR1) with Met at position 1. Carboxy terminal end
XX of recombinant RacOR1 has a covalently bound ligand binding moiety, which
XX can be a IL2 antibody directed against CD22 on cancerous B cells or human
XX chorionic gonadotropin (hCG) effective against Kaposi's sarcoma cells.
XX Recombinant ribonucleases can be expressed in bacteria without an N-
XX terminal methionine due to the presence of a signal peptide that is
XX cleaved by bacteria. The soluble expression of ribonuclease allows the
XX proteins to be fused in-frame with ligand binding moieties to form
XX cytotoxic fusion proteins. They can be used for treatment of cancer and
XX autoimmune diseases.
XX
XX Sequence 333 BP; 83 A; 95 C; 57 G; 98 T; 0 other:
XX
Query Match          100.0%; Score 330; DB 20; Length 333;
Best Local Similarity 100.0%; Pred. No. 4.4e-92;
Matches 330; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 cagaactgggtactcttcagcagaacacatcatcaaacctcgcgactcgtgaacact 60
DB 4 cagaactgggtactcttcagcagaacacatcatcaaacctcgcgactcgtgaacact 63
QY 61 atcattggacacacacatctacatcgtgtggtgcagtcgcaacagtggttaacatttc 120
DB 64 atcattggacacacacatctacatcgtgtggtgcagtcgcaacagtggttaacatttc 123
QY 121 atctctctgctactactgtttaagatcatctgactgtgtgtttaacaagaagttctg 180
DB 124 atctctctgctactactgtttaagatcatctgactgtgtgtttaacaagaagttctg 183
QY 181 tctactactcgtttccagctgaacactgtgactcgtactctatcaactccgctcgtgc 240
DB 184 tctactactcgtttccagctgaacactgtgactcgtactctatcaactccgctcgtgc 243
QY 241 cgtactctctcgtactcgtgaactaactacatctgcgtttaaagcgaaaacagttaccg 300
DB 244 cgtactctctcgtactcgtgaactaactacatctgcgtttaaagcgaaaacagttaccg 303
QY 301 gtccattcgtcgtgtatcgcgttgcccg 330

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CC terminal end of recombinant RacOR1 has a covalently bound ligand binding
CC moiety, which can be a L12 antibody directed against CD22 on cancerous B
CC cells or human chorionic gonadotrophin (hCG) effective against Kaposi's
CC sarcoma cells. Recombinant ribonucleases can be expressed in bacteria
CC without an N-terminal methionine due to the presence of a signal peptide
CC that is cleaved by bacteria. The soluble expression of ribonuclease
CC allows the proteins to be fused in-frame with ligand binding moieties to
CC form cytotoxic fusion proteins. They can be used for treatment of cancer
CC and autoimmune diseases.

Sequence 333 BP; 83 A; 95 C; 56 G; 99 T; 0 other;

Query Match 99.1%; Score 327; DB 20; Length 333;
Best Local Similarity 100.0%; Pred. No. 3.7e-91;
Matches 327; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 4 aactgggactcttcagcaggaacatatacatcaactccgactatctgcaacatc 63
DB 7 aactgggactcttcagcaggaacatatacatcaactccgactatctgcaacatc 66
OY 64 atgacaacacacatcatcagctgtgtgcaagtgtaaacattcatc 123
DB 67 atgacaacacacatcatcagctgtgtgcaagtgtaaacattcatc 126
OY 124 tctctgctactactgttaagctatctgcaagtgtaaacattcatc 183
DB 127 tctctgctactactgttaagctatctgcaagtgtaaacattcatc 186
OY 184 actactcgtttcagcaggaacatctgcaagtgtaaacattcatc 243
DB 187 actactcgtttcagcaggaacatctgcaagtgtaaacattcatc 246
OY 244 tactctctcgtactgtaaacatcatctgcaagtgtaaacattcatc 303
DB 247 tactctctcgtactgtaaacatcatctgcaagtgtaaacattcatc 306
OY 304 catttcgctgtatcgtgtgtgccc 330
DB 307 catttcgctgtatcgtgtgtgccc 333

RESULT 5

AA208132 ID AA208132 standard; cDNA; 330 BP.

AC AA208132;

DT 25-JAN-2000 (first entry)

DE Recombinant RacOR1 Met22Lau Met57Lau encoding cDNA.

XX Recombinant Rana catesbeiana oocyte ribonuclease; covalently bound;
KW RacOR1 Met22Lau Met57Lau; L12 antibody; ligand binding moiety; CD22;
KW cancerous B cell; Kaposi's sarcoma; human chorionic gonadotrophin; hCG;
KW signal peptide; recombinant ribonuclease; cytotoxic fusion protein;
KW cancer; bullfrog; autoimmune disease; RNase; ss.
XX Rana catesbeiana.
OS Synthetic.
XX

XX Key Location/Qualifiers

FT mat_peptide 1..330

FT /tag- a /product- "Recombinant RacOR1 Met22Lau Met57Lau"

FT /note- "Rana catesbeiana oocyte ribonuclease"

FT replace(64..66, ATG)

FT old_sequence /tag- b

FT old_sequence replace(169..171, ATG)

FT /tag- c

XX WO9950398-A2.

PD 07-OCT-1999.

XX 26-MAR-1999; 99WO-US06641.

XX 27-MAR-1998; 98US-0079751.

XX (USSH) US DEPT HEALTH & HUMAN SERVICES.

XX Newton DL, Rybak SM;

DR WPI: 1999-610847/52.

DR P-PSDB: AAY28874.

PT New recombinant ribonucleases, used for killing target cells, e.g. for

PS treating cancers, viral infections or autoimmune diseases

XX Disclosure; Page 64; 71pp; English.

CC The present sequence is a cDNA encoding recombinant Rana catesbeiana
CC oocyte ribonuclease (RacOR1) with Met22Lau Met57Lau. Carboxy terminal end
CC of recombinant RacOR1 has a covalently bound ligand binding moiety, which
CC can be a L12 antibody directed against CD22 on cancerous B cells or human
CC chorionic gonadotrophin (hCG) effective against Kaposi's sarcoma cells.
CC Recombinant ribonucleases can be expressed in bacteria without an N-
CC terminal methionine due to the presence of a signal peptide that is
CC cleaved by bacteria. The soluble expression of ribonuclease allows the
CC proteins to be fused in-frame with ligand binding moieties to form
CC cytotoxic fusion proteins. They can be used for treatment of cancer and
CC autoimmune diseases.

Sequence 330 BP; 80 A; 97 C; 56 G; 97 T; 0 other;

Query Match 99.0%; Score 326.8; DB 20; Length 330;
Best Local Similarity 99.4%; Pred. No. 4.3e-91;
Matches 328; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 1 cagaactgggactcttcagcaggaacatatacatcaactccgactatctgcaacact 60

DB 1 cagaactgggactcttcagcaggaacatatacatcaactccgactatctgcaacact 60

OY 61 atcatggacaacacatcatcagctgtgtgcaagtgtaaacattcatc 120

DB 61 atcatggacaacacatcatcagctgtgtgcaagtgtaaacattcatc 120

OY 121 atctctctgctactactgtaaacatcatctgcaagtgtaaacattcatc 180

DB 121 atctctctgctactactgtaaacatcatctgcaagtgtaaacattcatc 180

OY 181 tctactactcgtttccagcaggaacatctgcaagtgtaaacattcatc 240

DB 181 tctactactcgtttccagcaggaacatctgcaagtgtaaacattcatc 240

OY 241 cgttactctctgctactgtaaacatcatctgcaagtgtaaacattcatc 300

DB 241 cgttactctctgctactgtaaacatcatctgcaagtgtaaacattcatc 300

OY 301 gtccatttcgctgtatcgtgtgtgccc 330

DB 301 gtccatttcgctgtatcgtgtgtgccc 330

RESULT 6

AA208133 ID AA208133 standard; cDNA; 333 BP.

XX AA208133;

DT 25-JAN-2000 (first entry)

DE Recombinant Met(-1) RacOR1 Met22Lau Met57Lau encoding cDNA.

KW Recombinant Met(-1) Rana catesbeiana ribonuclease Met22Lau Met57Lau;

Query Match	Best Local Similarity	Matches	328; Conservative	0;	Mismatches	2;	Indels	0;	Gaps	0
QY	1	cagaactcgtgctactcttcacagcagaacaatcatcatcaatacattccgatactcgtcaaacact	60							
DB	4	cagaactcgtgctactcttcacagcagaacaatcatcatcaatacattccgatactcgtcaaacact	63							
QY	61	atcatcgcgaacaacatcatcgtctgtgtcagtcgaagcgttttaacattccatc	120							
DB	64	atctcgtgcaacaacatcatcgtctgtgtcagtcgaagcgttttaacattccatc	123							
QY	121	atctcttcgtcactactcgttlaaagctacatcgtcactcgtgtcttcaacaatgaacgtctc	180							
DB	124	atctcttcgtcactactcgttlaaagctacatcgtcactcgtgtcttcaacaactcgaacgtctc	183							
QY	181	tctactactcgttctcagctgaacactcgtcactcgttaactctatacctccgcgtctcgtc	240							

Query Match	28.8%; Score 95; DB 20; Length 95;
Best Local Similarity 100.0%; Pred. No. 9,6e-20;	
Matches 95; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	
QY 226 actcgcgctcgtgcgcgactctcttcgtactgaactaactacatctgcgttaaatgc 285	
Db 1 actcgcgctcgtgcgcgactctcttcgtactgaactaactacatctgcgttaaatgc 60	
QY 286 gaaacacgatacccggttcatttcgctgltacg 320	
Db 61 gaaacacgatacccggttcatttcgctgltacg 95	
RESULT 8	
AAZ08148	
ID AAZ08148 standard; DNA; 96 BP.	
XX	
CC AAZ08148;	
Db 184 tctactactcgttccagctgaacaactctgcactcgttactttatcaactccgcgctcgtgc 243	
QY 241 ccgactactcttcgttactgaactaactacatctcgtttaaattgcgaaacacgataccg 300	
Db 244 ccgactactcttcgttactgaactaactacatctcgtttaaattgcgaaacacgataccg 303	
QY 301 gttcaattcgcgtgtatcgcgttcgttgcccg 330	
Db 304 gttcaattcgcgtgtatcgcgttcgttgcccg 333	
RESULT 7	
AAZ08144	
ID AAZ08144 standard; DNA; 95 BP.	
XX	
AC AAZ08144;	
XX	
DT 25-JAN-2000 (first entry)	
XX	
DE PCR primer-3 for synthesising 3' half of RacOR1 gene.	
XX	
KW PCR primer: ribonuclease; RNase; RacOR1, Rana catesbeiana; mutation;	
KM recombinant RNase; ss.	
XX	
OS Synthetic.	
OS Rana catesbeiana.	
XX	
PN WO950398-A2.	
XX	
PD 07-OCT-1999.	
XX	
PE 26-MAR-1999; 99WO-US06641.	
XX	
PR 27-MAR-1998; 98US-0079751.	
XX	
PA (USSH) US DEPT HEALTH & HUMAN SERVICES.	
XX	
PI Newton DL, Rybak SM;	
XX	
DR WPI; 1999-610847/552.	
XX	
PT New recombinant ribonucleases, used for killing target cells, e.g. for	
XX	
PS treating cancers, viral infections or autoimmune diseases -	
XX	
PS Example 5; Page 41; 71pp; English.	
XX	
CC The present sequence is a PCR primer comprising the 3' half of	
CC ribonuclease (RacOR1) gene from Rana catesbeiana. It is used along with	
CC other primers to synthesise the 3' half of RNase with mutations resulting	
CC in recombinant RNase with Met32Leu and Met57Leu.	
XX	
SO Sequence 95 BP; 21 A; 29 C; 19 G; 26 T; 0 other;	

```
XX 25-JAN-2000 (first entry)
XX
XX PCR primer-2 for assembling mutated RacOR1 gene.
DE
XX PCR primer: assemble; Rana catesbelana ribonuclease gene; RacOR1; RNase;
XX mutated; ss.
XX
XX Synthetic.
OS Rana catesbelana.
XX
XX WO950398-A2.
XX
XX 07-OCT-1999.
XX
XX 26-MAR-1999; 99WO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
XX
XX Newton DL, Rybak SM;
XX
XX WPI; 1999-610847/52.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX Example 5; Page 41; 71pp; English.
XX
XX The present sequence is a PCR primer, used along with another primer
XX to assemble mutated Rana catesbelana ribonuclease (RacOR1) gene
XX resulting in RNase with Met22Leu and Met57Leu.
XX
XX Sequence 96 BP; 17 A; 32 C; 17 G; 30 T; 0 other;
SQ

Query Match      28.6%; Score 94.4; DB 20; Length 96;
Best Local Similarity 99.0%; Pred. No. 1.5e-19;
Matches 95; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 148 atctgcactggtgttatcaacatgaacgtctctctactactcgtttccagctgaacact 207
   |||||||
DB 1 atctgcactggtgttatcaacatgaacgtctctctactactcgtttccagctgaacact 60

OY 208 tgcactcgtactctctatcactcgcgcgtgcgcg 243
   |||||||
DB 61 tgcactcgtactctctatcactcgcgcgtgcgcg 96

RESULT 9
AAZ08141
ID AAZ08141 standard; DNA; 96 BP.
XX
XX AAZ08141;
AC
XX
XX 25-JAN-2000 (first entry)
DT
XX
XX PCR primer-1 for synthesizing 3' half of RacOR1 gene.
DE
XX
XX PCR primer: ribonuclease; RNase; RacOR1; Rana catesbelana; mutation;
XX recombinant RNase; ss.
XX
XX Synthetic.
OS Rana catesbelana.
XX
XX WO950398-A2.
XX
XX 07-OCT-1999.
XX
XX 26-MAR-1999; 99WO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
```

```
XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
XX
XX Newton DL, Rybak SM;
XX
XX WPI; 1999-610847/52.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX Example 5; Page 41; 71pp; English.
XX
XX The present sequence is a PCR primer comprising the 3' half of
XX ribonuclease (RacOR1) gene from Rana catesbelana. It is used along with
XX other primers to synthesise the 3' half of RNase with mutations resulting
XX in recombinant RNase with Met22Leu and Met57Leu.
XX
XX Sequence 96 BP; 18 A; 31 C; 17 G; 30 T; 0 other;
SQ

Query Match      28.1%; Score 92.8; DB 20; Length 96;
Best Local Similarity 97.9%; Pred. No. 4.6e-19;
Matches 94; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

OY 148 atctgcactggtgttatcaacatgaacgtctctctactactcgtttccagctgaacact 207
   |||||||
DB 1 atctgcactggtgttatcaacatgaacgtctctctactactcgtttccagctgaacact 60

OY 208 tgcactcgtactctctatcactcgcgcgtgcgcg 243
   |||||||
DB 61 tgcactcgtactctctatcactcgcgcgtgcgcg 96

RESULT 10
AAZ08147
ID AAZ08147 standard; DNA; 96 BP.
XX
XX AAZ08147;
AC
XX
XX 25-JAN-2000 (first entry)
DT
XX
XX PCR primer-1 for assembling mutated RacOR1 gene.
DE
XX
XX PCR primer: assemble; Rana catesbelana ribonuclease gene; RacOR1; RNase;
XX mutated; ss.
XX
XX Synthetic.
OS Rana catesbelana.
XX
XX WO950398-A2.
XX
XX 07-OCT-1999.
XX
XX 26-MAR-1999; 99WO-US06641.
XX
XX 27-MAR-1998; 98US-0079751.
XX
XX (USSH ) US DEPT HEALTH & HUMAN SERVICES.
XX
XX Newton DL, Rybak SM;
XX
XX WPI; 1999-610847/52.
XX
XX New recombinant ribonucleases, used for killing target cells, e.g. for
XX treating cancers, viral infections or autoimmune diseases -
XX
XX Example 5; Page 41; 71pp; English.
XX
XX The present sequence is a PCR primer, used along with another primer
XX to assemble mutated Rana catesbelana ribonuclease (RacOR1) gene
XX resulting in RNase with Met22Leu and Met57Leu.
XX
XX Sequence 96 BP; 30 A; 28 C; 16 G; 22 T; 0 other;
SQ
```

Query Match 27.6%; Score 91.2; DB 20; Length 96;
 Best Local Similarity 96.9%; Pred. No. 1.4e-18;
 Matches 93; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

OY 1 cagaactgggtactcttcagcagaatacatcatcaactcgcgtatctgcaacact 60
 |||||||
 DB 1 cagaactgggtactcttcagcagaatacatcatcaactcgcgtatctgcaacact 60
 |||||||

OY 61 atcatggacaacaacatcatcgttgggtgacag 96
 |||||
 DB 61 atccctgcagacaacatcatcgttgggtgacag 96
 |||||

RESULT 11
 AAZ08139
 ID AAZ08139 standard; DNA; 97 BP.
 AC AAZ08139;
 XX
 XX 25-JAN-2000 (first entry)
 DE PCR primer-1 for synthesizing 5' half of RACOR1 gene.
 XX
 XX PCR primer: ribonuclease; RNase; RACOR1; Rana catesbeiana; mutation;
 KW recombinant RNase; ss.
 XX
 XX Synthetic.
 OS Rana catesbeiana.
 XX
 XX Key Location/Qualifiers
 FH misC.feature 22
 FT /*tag" a
 FT /note- "Unknown additional base"
 XX
 XX W09950398-A2.
 PN 07-OCT-1999.
 PD
 XX 26-MAR-1999; 99WO-US06641.
 PF
 XX 27-MAR-1998; 98US-0079751.
 PR
 XX (USSH) US DEPT HEALTH & HUMAN SERVICES.
 PA
 XX Newton DL, Rybak SM;
 PI
 XX WPI: 1999-610847/52.
 DR
 XX New recombinant ribonucleases, used for killing target cells, e.g. for
 PT treating cancers, viral infections or autoimmune diseases -
 PT
 XX Example 5; Page 40; 71pp; English.
 PS
 XX The present sequence is a PCR primer comprising the 5' half of
 CC ribonuclease (RACOR1) gene from Rana catesbeiana. It is used along with
 CC other primers to synthesise the 5' half of RNase with mutations resulting
 CC in recombinant RNase with Met22Leu and Met57Leu.
 XX
 SQ Sequence 97 BP; 31 A; 27 C; 16 G; 22 T; 1 other;

Query Match 25.8%; Score 85; DB 20; Length 97;
 Best Local Similarity 99.0%; Pred. No. 1.2e-16;
 Matches 96; Conservative 0; Mismatches 0; Indels 1; Gaps 1;

OY 1 cagaactgggtactcttcagcagaatacatcatcaactcgcgtatctgcaaac 59
 |||||||
 DB 1 cagaactgggtactcttcagcagaatacatcatcaactcgcgtatctgcaaac 60
 |||||||

OY 60 tatcatggacaacaacatcatcgttgggtgacag 96
 |||||||

DB 61 tatcatggacaacaacatcatcgttgggtgacag 97
 |||||||

RESULT 12
 AAZ08140
 ID AAZ08140 standard; DNA; 86 BP.
 AC AAZ08140;
 XX
 XX 25-JAN-2000 (first entry)
 DE PCR primer-2 for synthesizing 5' half of RACOR1 gene.
 XX
 XX PCR primer: ribonuclease; RACOR1; Rana catesbeiana; mutation; RNase;
 KW recombinant RNase; ss.
 XX
 XX Synthetic.
 OS Rana catesbeiana.
 XX
 XX W09950398-A2.
 PN 07-OCT-1999.
 PD
 XX 26-MAR-1999; 99WO-US06641.
 PF
 XX 27-MAR-1998; 98US-0079751.
 PR
 XX (USSH) US DEPT HEALTH & HUMAN SERVICES.
 PA
 XX Newton DL, Rybak SM;
 PI
 XX WPI: 1999-610847/52.
 DR
 XX New recombinant ribonucleases, used for killing target cells, e.g. for
 PT treating cancers, viral infections or autoimmune diseases -
 PT
 XX Example 5; Page 40; 71pp; English.
 PS
 XX The present sequence is a PCR primer comprising the 5' half of
 CC ribonuclease (RACOR1) gene from Rana catesbeiana. It is used along with
 CC other primers to synthesise the 5' half of RNase with mutations resulting
 CC in recombinant RNase with Met22Leu and Met57Leu.
 XX
 SQ Sequence 86 BP; 19 A; 20 C; 16 G; 31 T; 0 other;

Query Match 21.8%; Score 71.8; DB 20; Length 86;
 Best Local Similarity 96.6%; Pred. No. 1.3e-12;
 Matches 84; Conservative 0; Mismatches 2; Indels 1; Gaps 1;

OY 79 tacatcgttgggtgcagcagaacgtgttaacacttcatcatctctctgtactact 138
 |||||||
 DB 1 tacatcgttgggtgcagcagaacgtgttaacacttcatcatctctctgtactact 59
 |||||||

OY 139 gttaaagctatctgcactggtgttattc 165
 |||||||
 DB 60 gttaaagctatctgcactggtgttattc 86
 |||||||

RESULT 13
 AAZ19767
 ID AAZ19767 standard; CDNA; 318 BP.
 AC AAZ19767;
 XX
 XX 01-DEC-1999 (first entry)
 DE Recombinant frog Onconase CDNA.
 XX
 XX Ribonuclease; protein synthesis; inhibition; cancer; cytotoxic; ds.
 KW
 XX Rana pipiens.
 OS
 XX

FH	Key	Location/Qualifiers
FT	CDS	1..318
FF		/tag=^a
FF		/product="Recombinant frog Onconase"
xx		
PN	WO9946389-A1.	
XX		
PD	16-SEP-1999.	
XX		
Pf	11-MAR-1999;	99WO-US04252.
XX		
PR	11-MAR-1998;	98US-0077557.
XX		
PA	(IMMU-) IMMUNOMEDICS INC.	
DR	Goldenberg DM, Hansen H, Leung S;	
PI	WPI; 1999-551416/46.	
DR	P-PsDB; AAI39400.	
XX		
PT	A new recombinant Onconase used to treat, e.g. colon cancer -	
PS	Example 1; Fig 1; 42pp; English.	
XX		
CC	This sequence represents recombinant frog Onconase cDNA. Onconase has	
CC	ribonuclease and anti-tumour activity. The cDNA was produced via PCR	
CC	(using primers AAZ19768-219769) of two synthetic DNAs whose sequences	
CC	encoded most of the N-terminal or the C-terminal amino acids of mature	
CC	Onconase. The two PCR products generated encoded either the N-terminal	
CC	54 amino acids (minus the initial methionine) or the C-terminal 51 amino	
CC	acids, and were ligated in frame at an NruI site. The cDNA was then	
CC	subcloned into a vector e.g., pBluescript, where the ATG initiation	
CC	codon was ligated to the cDNA. After expression in E. coli, the	
CC	recombinant protein was purified. The initial N-formyl methionine was	
CC	cleaved off and the now N-terminal glutamate residue cyclised to form an	
CC	N-terminal pyroglutamate. The pyroglutamate residue forms part of the	
CC	phosphate binding pocket of Onconase and is essential for both	
CC	ribonuclease and anti-tumour activity. Onconase is a 12 kD ribonuclease	
CC	which causes cell death as a result of potent inhibition of protein	
CC	synthesis by a mechanism involving inactivation of cellular RNA. It is	
CC	not inhibited by mammalian placental ribonuclease inhibitor, which may	
CC	explain its enhanced cytotoxicity relative to mammalian enzymes. It has	
CC	anti-tumour activity against a variety of solid tumours e.g., colon or	
CC	pancreatic cancers, and can be used alone or in combination with other	
CC	anti-cancer agents such as tamoxifen. When used as an anti-tumour agent,	
CC	Onconase can be conjugated to a marker which targets it to a specific	
CC	cell type.	
SQ	Sequence 318 BP; 99 A; 65 C; 71 G; 83 T; 0 other;	
Query Match	15.2%; Score 50; DB 20; Length 318;	
Best Local Similarity	54.2%; Pred. No. 1.2e-05;	
Matches 179; Conservative	0; Mismatches 130; Indels 21; Gaps	3
OY	1 cagaactgggctacttccagcagaacaatcatcaaacatcgatactgcgaacct 60 Db 4 caggattggctaagcttcagaagaacatcacgaataacc-----gagatgtga 54	
OY	61 atcatgtaacaacaacatcaacatcgtttggtgcagtgcgaacggttaaacattatc 120 Db 55 gactgcgaacaatatatgtctacgatctggtttcactgtaaagataagataaccttata 114	
OY	121 acctcttcgtactacatgtttaagctatcgactcgatggttatc--aacatgaagt 177 Db 115 tecaagtggcgagagccvtgtaaaggctacvtytaaggatatalccgagtaagaacgtg 174	
OY	178 ctgtctactactcgtttccaagcttgtaaacattgcactctatactaactcgcgtcg 237 Db 175 cgtactactccgaggttctatctgtccagttgcaatgtgactta-----cggcc 225	
OY	238 tgcgcgtactcttctcgtaataactaatcatctgcttaaatgcgaaaacgaactac 297 	

Db	226	lqcaaatlaagctgtaagaagaaagactaacaaatttgcgtaactlqcgagaccagct	285
Oy	298	ccggatcatccgcgtgatacgatgcgtgtgc	327
Db	286	ccgtacatcttcgttgtagtgcggagagctgc	315
RESULT 14			
ID	AAS34685		
XX	AAS34685 standard; DNA; 19205 BP.		
AC	AAS34685:		
XX			
DT	17-DEC-2001 (first entry)		
XX			
DE	Human DNA for a novel foetal antigen, SEQ ID No 2109.		
KW	Human; foetal tissue antigen; ds; antiinflammatory; neuroprotective;		
KW	immunomodulator; cardiovascular; cytosolic; nephrotoxic;		
KW	cardiovascular; autoimmune disease; rheumatoid arthritis;		
KW	hyperproliferative disorder; breast neoplasm; cancer;		
KW	cardiovascular disorder; cardiac arrest; cerebrovascular disorder;		
KW	cerebral ischaemia; angiogenesis; nervous system disorder;		
KW	Alzheimer's disease; infection; ocular disorder; corneal infection;		
KW	wound healing; epithelial cell proliferation; food additive.		
XX			
OS	Homo sapiens.		
XX			
PM	WO200155312-A2.		
XX			
PD	02-AUG-2001.		
XX			
PF	17-JAN-2001; 2001WO-US01321.		
XX			
PR	31-JAN-2000; 2000US-0179065.		
PR	04-FEB-2000; 2000US-0180628.		
PR	24-FEB-2000; 2000US-0184664.		
PR	02-MAR-2000; 2000US-0186350.		
PR	16-MAR-2000; 2000US-0189874.		
PR	17-MAR-2000; 2000US-0190076.		
PR	18-APR-2000; 2000US-0198123.		
PR	19-MAY-2000; 2000US-0205515.		
PR	07-JUN-2000; 2000US-0209467.		
PR	28-JUN-2000; 2000US-0214886.		
PR	30-JUN-2000; 2000US-0215135.		
PR	07-JUL-2000; 2000US-0216647.		
PR	07-JUL-2000; 2000US-0216880.		
PR	11-JUL-2000; 2000US-0217487.		
PR	11-JUL-2000; 2000US-0217496.		
PR	14-JUL-2000; 2000US-0218290.		
PR	26-JUL-2000; 2000US-0220963.		
PR	26-JUL-2000; 2000US-0220964.		
PR	14-AUG-2000; 2000US-0224518.		
PR	14-AUG-2000; 2000US-0224519.		
PR	14-AUG-2000; 2000US-0225213.		
PR	14-AUG-2000; 2000US-0225214.		
PR	14-AUG-2000; 2000US-0225266.		
PR	14-AUG-2000; 2000US-0225267.		
PR	14-AUG-2000; 2000US-0225268.		
PR	14-AUG-2000; 2000US-0225270.		
PR	14-AUG-2000; 2000US-0225447.		
PR	14-AUG-2000; 2000US-0225757.		
PR	14-AUG-2000; 2000US-0225758.		
PR	14-AUG-2000; 2000US-0225759.		
PR	18-AUG-2000; 2000US-0226279.		
PR	22-AUG-2000; 2000US-0226681.		
PR	22-AUG-2000; 2000US-0226868.		
PR	22-AUG-2000; 2000US-0227182.		
PR	23-AUG-2000; 2000US-0227009.		
PR	30-AUG-2000; 2000US-0228924.		
PR	01-SEP-2000; 2000US-0229287.		
PR	01-SEP-2000; 2000US-0229343.		
PR	01-SEP-2000; 2000US-0229344.		


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Db      1017 ctaattatcatatcatcattcgatccaacagtttctcccttaaccccttagaccctcaca 1076
QY      220 tctatcacctcgcgcgtgcgcgtacctctcttcgtactgaaactaactacatcgtgct 279
        ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      1077 cctaacctaccctctccccaataactactattatatttgaaaattttactgctctttc 1136
QY      280 aaatgcga 287
        || |||||
Db      1137 aactgcga 1144

RESULT 15
ABA44186
ID      ABA44186 standard; DNA; 436 BP.
XX
AC      ABA44186;
XX
DT      01-FEB-2002 (first entry)
XX
DE      Human breast cell single exon nucleic acid probe #2881.
XX
KW      Human; microarray; single exon probe; gene expression; breast;
KM      disease; cancer; ss.
XX
OS      Homo sapiens.
XX
PN      MO200157271-A2.
XX
PD      09-AUG-2001.
XX
PE      30-JAN-2001; 2001MO-US000662.
XX
PR      04-FEB-2000; 2000US-0180312.
PR      26-MAY-2000; 2000US-0207456.
PR      30-JUN-2000; 2000US-0608408.
PR      03-AUG-2000; 2000US-0632366.
PR      21-SEP-2000; 2000US-0234687.
PR      27-SEP-2000; 2000US-0236359.
PR      04-OCT-2000; 2000GB-0024263.
XX
PA      (MOLE-) MOLECULAR DYNAMICS INC.
XX
PI      Penn SG, Hanzel DK, Chen W, Rank DR;
XX
DR      MPI; 2001-496933/54.
XX
PT      New spatially-addressable set of single exon nucleic acid probes,
PT      useful for measuring gene expression in sample derived from human
PT      breast, comprises number of single exon nucleic acid probes -
XX
PS      Claim 1; SEQ ID NO 2881; 327bp + sequence listing; English.
XX
CC      The invention relates to a spatially-addressable set of single exon
CC      nucleic acid probes for measuring gene expression in a sample derived
CC      from human breast and Br 474 cells. The method involves contacting
CC      the probes with a collection of detectably labelled nucleic acids
CC      derived from mRNA of human breast, and then measuring the label
CC      bound to each probe of the microarray. The probes are useful for
CC      verifying the expression of regions of genomic DNA predicted to
CC      encode proteins. They are useful for gene discovery and for
CC      determining predisposition and/or prognosing breast disease. Gene
CC      expression analysis is useful for assessing the toxicity of chemical
CC      agents on cells. The microarray of this invention presents a far greater
CC      diversity of probes for measuring gene expression, with far less bias
CC      than expressed sequence tag microarrays. The method is suitable for
CC      rapid production of functional information from genomic sequence. The
CC      present sequence is a single exon nucleic acid probe of the invention.
CC      Note: The sequence data for this patent did not form part of the
CC      printed specification, but was obtained in electronic format directly
CC      from WIPO at ftp.wipo.int/pub/published_pct_sequences.
XX
SQ      Sequence 436 BP; 137 A; 144 C; 15 G; 140 T; 0 other;

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Query Match      11.8%; Score 39; DB 22; Length 436;
Best Local Similarity 47.4%; Pred. No. 0.035;
Matches 117; Conservative 0; Mismatches 130; Indels 0; Gaps 0;

QY      34 atcaacacatccgatcatctgcgaacactatcatatggaacaacaacatcactcgttgyt 93
        ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      112 atcactactaccactaccactactactactactactactactactactactactaccgact 171
QY      94 cagtgcaaaagtggttaacactttcatcatctctcgtctactcgttccagctgaacacttgctc 153
        ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      172 atcattactactactaccacccatccatcatcactactactactactactactactactact 231
QY      154 actggtgtatcaacaacatgcagttctgtcactactcgtttccagctgaacacttgact 213
        || | ||| | ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      232 gcttctataattactactcctactcctaccattactactactactactactactactactact 291
QY      214 cgtacttctatcactccggtcgcgtgcgtactctctcgtactgaactaactatcact 273
        ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db      292 actactactatcatlactactataccacacatcactactactactactactactactaccacc 351
QY      274 tgcgtta 280
        || |||
Db      352 accatla 358

```

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